

GenCore version 4.5														
Copyright (c) 1993 - 2000 Compugen Ltd.														
nucleic - nucleic search, using sw model														
on : March 15, 2002, 03:41:53 ; Search time 1909.32 Seconds (without alignments) 24735.349 Million cell updates/sec														
title:	US-09-652-292-1	strict score:	4395	sequence:	1 gagggggtccctgtccaggcc.....atttttgtaaaaaaa 4395	scoring table:	OLIGO_NUC	gapop 60.0 , Gapext 60.0	searched:					
third size :	0	total number of hits satisfying chosen parameters:	11351937 seqs, 5372889281 residues	minimum DB seq length:	0	maximum DB seq length:	2000000000	start-processing: Listing first 45 summaries	database :					
EST:*	1: em_estfun:*	2: em_esthum:*	3: em_estin:*	4: em_estom:*	5: em_estpl:*	6: em_estba:*	7: em_estro:*	8: em_estov:*	9: em_nhcc:*					
	10: qb_est1:*	11: qb_est2:*	12: qb_itrc:*	13: qb_gss:*	14: em_gss_fun:*	15: em_gss_hum:*	16: em_gss_inv:*	17: em_gss_pln:*	18: em_gss_pro:*					
	19: em_gss_rdt:*	20: em_gss_vrt:*	21: em_gss_other:*											
Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.														
SUMMARIES														
result:	AQ037826	locus:	CIT-HSP-2326P11	definition:	CIT-HSP-2326P11.TF	sequence:	GSS	11-JUL-1998						
	AQ037826								genomic clone 2326P11, DNA					
organism:	Homo sapiens	source:	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.	reference:	1 (bases 1 to 678)	authors:	Adams, M.D., Rounseley, S.D., Zhao, S., Field, C.E., Bass, S., Linher, K., Golden, K., Berry, K., Granger, D., Sub, E., Wible, C., Shizuya, H., Simon, M. and Ventner, J.C.	title:	Use of a random BAC End Sequence Database for Sequence-ready Map Building (1998)					
comment:	Other_GSS: CIT-HSP-2326P11.TR	contact:	Mark Adams	department:	The Institute for Genomic Research	address:	9712 Medical Center Dr., Rockville, MD 20850, USA	tel:	301 838 0200					
									Fax: 301 838 0208					
									Email: mdadams@igb.org					
									Clones are available from Research Genetics (http://www.regen.com). BAC end search page: http://www.tigr.org/tgb/humgen/bac_end_search/bac_end_search.html . Seq primer: M13-21					
									Class: BAC ends					
									features: Location/Qualifiers					
									/organism="Homo sapiens"					
									1. F78					
									2. 859 11					
									BG563179 602584689					
									3. 384 8.7					
									BG563179 602584689					

Search completed: March 15, 2002, 05:44:10